

## **Dengue Hemorrhagic Fever Virus in Saudi Arabia: A Review**

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**Abstract:**

Dengue fever is a global disease with a spectrum of clinical manifestation ranging from mild febrile disease to a severe disease in the form of dengue hemorrhagic fever and dengue shock syndrome. Dengue virus is one viral hemorrhagic fever that exists in the Kingdom of Saudi Arabia in addition to Alkhurma (Alkhurma) Hemorrhagic Fever, Chikungunya virus, Crimean–Congo Hemorrhagic Fever, and Rift Valley Fever. The disease is limited to the Western and South-western regions of Saudi Arabia where *Aedes aegypti* exists. The majority of the cases in Saudi Arabia had mild disease and is related to serotypes 1-3 but not 4. The prospect for Dengue virus control relies on vector control, health education and possibly vaccine use. Despite extensive collaborative efforts between multiple governmental sectors including Ministry of Health, Ministry of Municipalities and Rural Affairs, and Ministry of Water dengue remains a major public health concern in the regions affected.

**Keywords:**

Dengue hemorrhagic fever virus; DHFV;

## **Introduction:**

Dengue fever (DF) is a global disease with a spectrum of clinical manifestation ranging from mild febrile disease to a severe disease in the form of dengue hemorrhagic fever (DHF) and dengue shock syndrome (DSS). According to the World Health Organization, severe dengue disease in suspected dengue patients is defined as the presence of any of severe plasma leakage that leads to shock (dengue shock) and/or fluid accumulation with respiratory distress; severe bleeding; or severe organ impairment (World Health Organization (WHO) 2012).

Dengue infection occurs in an endemic form in 128 countries worldwide (Khetarpal & Khanna 2016). DF is one of several viral hemorrhagic fevers that exist in the Kingdom of Saudi Arabia (Alhaeli et al. 2016) in addition to Alkhurma (Alkhurma) Hemorrhagic Fever (AHF) (Zaki 1997; Al-Tawfiq & Memish 2017), Chikungunya Haemorrhagic Fever (Hussain et al. 2013), Crimean–Congo Hemorrhagic Fever (CCHF) (Leblebicioglu et al. 2015; El-Azazy & Scrimgeour 1997; Hassanein et al. 1997), and Rift Valley Fever (RVF) (Balkhy & Memish 2003; Al-Afaleq & Hussein 2011). Of the four serotypes of Dengue virus, serotypes 1-3 but not 4 were reported in Saudi Arabia (Ashshi 2017). The first description of Dengue fever in Saudi Arabia dates back to 1994 when Dengue virus serotype 2 (DEN-2) was isolated from a fatal and a non-fatal cases in Jeddah, Saudi Arabia (Fakeeh & Zaki 2001).

## **Search strategy**

The search included MEDLINE and Scopus databases for articles published in English as follows:

#1: “Dengue” OR “Dengue Virus” OR “Dengue Hemorrhagic Fever” OR “Dengue Fever”

#2: “Saudi Arabia” OR “Kingdom of Saudi Arabia”;

#3: #1 AND #2.

In addition, we searched the Saudi Epidemiology Bulletin (available from the Saudi Ministry of Health at: <http://www.fetp.edu.sa/Bulletin.html> ).

### **The Virus:**

Dengue fever, DHF, and DSS are caused by the dengue viruses (DENV). Dengue virus is a member of the genus *Flavivirus*, a member of the family *Flaviviridae*. The genus *Flavivirus* is classified into two broad categories based on the vector of transmission: tick-borne viruses and mosquito-borne viruses (figure 1). The Dengue virus is a mosquito-borne virus. Four antigenically related but distinct dengue virus serotypes exist and are: dengue virus types 1–4 (DENV-1, DENV-2, DENV-3, and DENV-4) (Khetarpal & Khanna 2016).

### **Virus Transmission:**

The Dengue viruses are transmitted to humans by the females of the mosquito *Aedes*. The most important vector is *A. aegypti* and other species such as *A. albopictus*, *A. polynesiensis*, and *A. niveushave* are secondary vectors (Malavige et al. 2004). The virus is limited to the Western and South-western regions of Saudi Arabia (Fakeeh & Zaki 2001; El-Kafrawy et al. 2016; Ayyub et al. 2006; Khan et al. 2008; Al-Azraqi et al. 2013; Alhaeli et al. 2016). Four different *Aedes species* were identified in the Western part of Saudi Arabia (Alikhan et al. 2014; Abdullah & Merdan 1995; Jupp et al. 2002; Al Ahmad et al. 2011; Alahmed et al. 2009; Kheir et al. 2010; Aziz et al. 2012) and only *A. caspius* was identified in the Eastern Saudi Arabia (Wills et al. 1985) (table 1). In a case control study, the following factors were associated with the risk of

Dengue virus infection: presence of stagnant water (OR = 4.9), indoor larvae (OR = 2.2), construction sites (OR = 2.2), and older age (OR = 1.2) (Kholed et al. 2012). It is known that rain fall in Jeddah is low. The occurrence of *A. aegypti* with Dengue fever in Jeddah is in paradox with the low level of rain; however, water containers play a role as breeding sites for *A. aegypti* (Ghaznawi et al. 1997; El-Gilany et al. 2010). In addition, the occurrence of huge constructions between 2008 and 2012 in Makkah, Saudi Arabia may had resulted in increased number of cases due to the formation of stagnant water (alwafi et al. 2013). Moreover, Jeddah is the Hajj entry point and is the largest commercial port and airport welcoming many Pilgrims coming from Dengue High Disease Burden. Thus in this context, the role of International Travel as the source of Dengue is a possibility. A recent study showed the role of visitors from dengue endemic countries in the importation of the virus into Saudi Arabia (Al-Saeed et al. 2017). The study showed that all dengue viruses in 2010-2015 were from the circulating Indian subcontinent lineage of the Cosmopolitan genotype (Al-Saeed et al. 2017).

### **Incidence:**

In Saudi Arabia, Dengue fever registry was made electronically in 2008 and Dengue is a notifiable disease in Saudi Arabia. The incidence of DENV infection among tested patients varies based on the location, year of the study and the method of testing (table 2) (Ashshi 2017; Ashshi et al. 2017; Fakeeh & Zaki 2001; Khan et al. 2008; Al-Azraqi et al. 2013; Ayyub et al. 2006; Organji et al. 2017; Shahin et al. 2009; Fakeeh & Zaki 2003; El-Gilany et al. 2010; alwafi et al. 2013; Gamil et al. 2014; Memish et al. 2011). The number of DENV infection was 6512 cases in 2013; 2081 cases in 2014; 4312 cases in 2015. The number of cases varies between 425 and 4312 per year (Alshamrani et al. n.d.; Organji et al. 2017). The annual number of cases of Dengue virus in Saudi Arabia is shown in figure 2 and the highest number of cases was in

2013, 2015, and 2016. However, the overall prevalence of DENV is 40-48.7% among clinically suspected patients (Khan et al. 2008; Ayyub et al. 2006) and 31.7% among random sample of patients attending the outpatients' clinics (Al-Azraqi et al. 2013). In a study from 2008 to 2012, the incidence rate doubled to 110 per 100,000 population in 2009 indicating the occurrence of an outbreak (alwafi et al. 2013). The majority of affected patients are adults and infected children constituted 24% (Shahin et al. 2009) in one study and 6% in another study (Ayyub et al. 2006). In a recent study, the age-standardised incidence rates of dengue was 10-99 per 100 000 person-years in 2013 (Stanaway et al. 2016).

### **Seasonality**

In a study of 159 cases in Makkah, 77% of the cases were during the spring and early summer (Shahin et al. 2009). And another study showed increased cases in the summer months and during the months of December and January (Kholed et al. 2012). In a study of 4187 cases, the peak cases occurred in April-May (alwafi et al. 2013) and a similar finding was in a report of 264 cases from Jazan (Gamil et al. 2014) and a study of cases in 2013-2014 (Aziz, Salman Abdo Al-Shami, et al. 2014). Thus, the majority of cases occurred in April-May.

### **Clinical Presentations:**

Dengue fever is characterized by constitutional findings of fever, severe headache, backache, joint pains, nausea and vomiting, eye pain and rash. The disease affects all age groups but tends to cause milder disease in young children. Dengue virus may cause one of four syndromes/diseases: undifferentiated fever, classic dengue fever, dengue hemorrhagic fever, or dengue shock syndrome. The first 207 patients had mild Dengue fever and only one patient had dengue shock syndrome (DSS) and one had Dengue Hemorrhagic fever (DHF) (Fakeeh & Zaki

2001). The affected patients in Saudi Arabia were more likely to be male and of young age group (summarized in table 3). The signs of symptoms of Saudi patients with dengue infections are summarized in table 4 (Ayyub et al. 2006; Khan et al. 2008; El-Gilany et al. 2010; Shahin et al. 2009; Ahmed 2010; Badreddine et al. 2017). The majority of patients (60-93%) presented with dengue fever, 5-39.4% had DHF, and about 1% had DSS. The reported mortality was also low. Dengue infection accounts for a total of 0.15-0.29 mortality per million person-years in Saudi Arabia in 2013 (Stanaway et al. 2016). Men are more affected than women in the various included studies. This is mainly related to the fact that men work outdoors and that women in Saudi wear clothing covering head to toes (alwafi et al. 2013).

### **Geographic Distribution**

Dengue virus was mainly reported from the Western and South-western regions of Saudi Arabia (Fakeeh & Zaki 2001; El-Kafrawy et al. 2016; Ayyub et al. 2006; Khan et al. 2008; Al-Azraqi et al. 2013; Alhaeli et al. 2016). This geographic restriction is directly related to the presence of *A. aegypti* in the region (Alikhan et al. 2014; Jupp et al. 2002; Al Ahmad et al. 2011; Alahmed et al. 2009; Kheir et al. 2010; Aziz et al. 2012) and not in other parts of the Kingdom of Saudi Arabia (Wills et al. 1985; Abdullah & Merdan 1995). Mathematical modelling showed that central Jeddah districts were the hotspots and the pattern changes greatly with time (Khormi et al. 2011). Using modelling techniques, a total of 111 districts in Jeddah were investigated for the risk of Dengue fever (Khormi & Kumar 2012). Of those districts, 15% were high risk, 22% were medium risk, 16% were low risk and 46% were very low risk (Khormi & Kumar 2012). An analysis of 2288 cases of Dengue fever, the disease was found to be concentrated in the south and central-north regions of Jeddah, Saudi Arabia (Alzahrani et al. 2013).

**Virus Serotypes:**

Dengue virus serotype is associated with the risk of DHF with highest risks with DENV-2, DENV-3, DENV-4 and DENV-1, as well as the pre-existence of anti-dengue antibodies. In the initial study of 985 suspected cases, DEN-2 accounted for 138 (66.7%) of 207 isolates, DEN-1 for 56 (27%), and DEN-3 for 13 (6.3%) (Fakeeh & Zaki 2001). The contribution of each serotype to Dengue in Saudi Arabia is shown in table 3. However, DENV-4 was not reported in any of the studies based on serology and molecular testing (Fakeeh & Zaki 2001; Ayyub et al. 2006; Khan et al. 2008; Organji et al. 2017; Fakeeh & Zaki 2003). Phylogenetic analysis of 19 isolates showed that DENV-1 and DENV-2 caused the 1994 outbreaks and it was an America-Africa genotype (lineage India-2) (Zaki et al. 2008). DENV-3 was isolated in 1997 and the outbreak in 2005-2006 was caused by a strain from genotype Asia (lineage Asia-2) (Zaki et al. 2008). Sequencing of the Dengue virus DENV-1-Jeddah-1-2011 strain showed high similarity with the Asian genotype (D1/H/IMTSSA/98/606 isolate) reported from Djibouti in 1998 (Azhar et al. 2015).

**Seroprevalence among asymptomatic individuals:**

The seroprevalence of Dengue virus antibodies among asymptomatic individuals was found to be 47.8% (927/1939) and among blood donors was 37% (68/184) (Jamjoom et al. 2016). The seroprevalence of anti-dengue IgG was 31.7% among asymptomatic persons attending outpatient clinics (Al-Azraqi et al. 2013). In one study, male gender, older age and communal and multi-story housing were significant factors for positive ELISA tests (Jamjoom et al. 2016). In a seroprevalence study of 1024 soldiers, only 0.1% tested positive for DENV by ELISA (Memish et al. 2011). Thus, there is variable seroprevalence of dengue among the different population



studied and is higher among patients attending outpatient clinics (31.7%) than the general population (0.1%). The general population may also represent the different regions of the country which may not be dengue fever areas.

### **Prospect for Control:**

Strategies to control Dengue virus requires the control of the vector, *A. aegypti*, through elimination of breeding sites and the elimination of the vector itself. It is important to intensify the use of insecticides to control mosquito due to the quick and efficient knock-down activity (Aziz, Salman Abdo Al-Shami, et al. 2014). In one study in Jazan, Saudi Arabia, *A. aegypti* mosquitoes were susceptible to Cyfluthrin and had variable resistances to other insecticides such as: lambda-cyhalothrin, Deltamethrin, Permethrin, Fenitrothion, Bendiocarb and DDT (Alsheikh et al. 2016). Health education and awareness of the disease and its vector play a major role in the control of Dengue in Saudi Arabia (Aziz, Salman A Al-Shami, et al. 2014). In one study from Saudi Arabia, high students' knowledge score was associated with family history of Dengue fever, having literate mothers, and age  $\geq 17$  years (Ibrahim et al. 2009). *Gambusia holbrooki* fish was effective in domestic water containers to control *A. aegypti* (Gamal 2012). In addition, the World Health Organization provides 36 boxes for the control of Dengue fever and includes *Aedes* control methods, Global Strategy for prevention and control of DF/DHF, and lessons learned from sustained efforts in countries combating dengue virus (Parks & Lloyd 2004). One dengue virus vaccine was licensed in Latin America and Southeast Asia. Two large phase III randomized controlled trials of this vaccine showed about 60% against virologically confirmed dengue in the first 13 months post-vaccine (Villar et al. 2015; Capeding et al. 2014). In a meta-analysis of nine studies, the vaccine efficacy was 54% with reduced efficacy of 34% for DENV2 (Malisheni et al. 2017). However, the vaccine was associated with higher relative risk of dengue

infection during the third year post-vaccination (Hadinegoro et al. 2015). The World Health Organization does not recommend the use of the vaccine for widespread vaccination nor for the use in areas with less than 50% seroprevalence (Anon 2016; World Health Organization 2017).

### **Conclusion:**

Dengue fever in Saudi Arabia is limited to the Western and South-western regions of the country and is linked to *Aedes aegypti*. The majority of the patients had mild disease and were caused by serotypes 1-3. Despite extensive collaborative efforts between multiple governmental sectors including Ministry of Health, Ministry of Municipalities and Rural Affairs, and Ministry of Water, dengue remains a major public health concern in the regions affected. The prospect for Dengue virus control relies on vector control, health education and possibly vaccine use.

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**Table 1: *Aedes* species according to the Region in Saudi Arabia**

<b>Species</b>	<b>Region</b>	<b>Year</b>	<b>Reference</b>
<i>Aedes aegypti</i> , <i>Aedes arabiensis</i> and <i>Aedes caspius</i> .	Western	1956	(Alikhan et al. 2014)
<i>Aedes caspius</i>	Eastern	1985	(Wills et al. 1985)
<i>Aedes caspius</i>	South western	1995	(Abdullah & Merdan 1995)
<i>Aedes vittatus</i>	Southern	2001	(Alikhan et al. 2014)
<i>Aedes vexans</i> , <i>Aedes arabiensis</i> , <i>Aedes vittatus</i> , <i>Aedes caspius</i> and <i>Aedes caballus</i>	Jizan	2002	(Jupp et al. 2002)
<i>Aedes caspius</i> and <i>Aedes aegypti</i>	Jeddah	2011	(Al Ahmad et al. 2011)
<i>Aedes aegypti</i> , <i>Aedes (Ochlerotatus) caspius</i> , <i>Aedes (Ochlerotatus) vexans</i> var. <i>arabiensis</i>	Jeddah	2014	(Alikhan et al. 2014)

<i>Aedes caspius</i> , <i>A. aegypti</i> and others	Makkah	2004-2006	(Alahmed et al. 2009)
<i>Aedes caspius</i> , <i>A. aegypti</i> and others	Madinah	2010	(Kheir et al. 2010)
co-breeding of <i>Aedes</i> , <i>Culex</i> and <i>Anopheles</i>	Makkah	2008-2009	(Aziz et al. 2012)

**Table 2: A Summary of the incidence of Dengue virus infection among tested patients based on the location, year of the study and the method of testing**

<b>Study type</b>	<b>Study population</b>	<b>Method of detection</b>	<b>Number included</b>	<b>% positive</b>	<b>Reference</b>
Cross sectional	Male blood donors	ELISA	910	39 IgG; 5.5 IgM	(Ashshi 2017; Ashshi et al. 2017)
Longitudinal	Suspected cases	Viral culture	985	21	(Fakeeh & Zaki 2001)
Longitudinal	Suspected cases	ELISA	985	11	(Fakeeh & Zaki 2001)
Cross sectional	random sample of patients attending the outpatients'	ELISA	965	31.7 IgG	(Al-Azraqi et al. 2013)



	clinics in Jizan and Aseer region				
Longitudinal April to July 2004	clinically suspected patients, Makkah	ELISA and RT- PCR	160	40 (n=64) IgM ELISA, 8.7 (n=14) by RT- PCR and 8.1 (n =13) by both	(Khan et al. 2008)
Longitudinal May 2004- April 2005	clinically suspected patients, Jeddah	ELISA	80	48.7	(Ayyub et al. 2006)
NA	clinically suspected patients, Makkah	RT-PCR	25	24	(Organji et al. 2017)
Longitudinal 2006 to 2008	clinically suspected patients, Makkah	ELISA or RT- PCR	159	100	(Shahin et al. 2009)

Longitudinal 1994 to 2002	clinically suspected patients, Jeddah	Virus isolation or ELISA	1020	31.3 (of those 65.5% by virus isolation and the rest were based on serology)	(Fakeeh & Zaki 2003)
September to mid December in 2006	Admitted patients, Makkah	Virus isolation or ELISA	71	100	(El-Gilany et al. 2010)
Longitudinal study 2008- 2012, Makkah	Confirmed cases	NA	4187	100	(alwafi et al. 2013)
Cross sectional April 2010- March 2011, Jazan	Suspected cases	ELISA	553	47.7	(Gamil et al. 2014)
Cross sectional,	Seroprevalence	ELISA	1024	0.1	(Memish et al.

2009, Saudi military forces, Jazan					2011)
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**Table 3: Summary of Characteristics of Confirmed Dengue Fever Cases in Saudi Arabia**

<b>Number</b>	<b>Study population</b>	<b>Method of detection</b>	<b>Serotype</b>	<b>Age</b>	<b>Male to Female ratio</b>	<b>Reference</b>
91	clinically suspected patients, Makkah	ELISA and RT-PCR	DENV-2: 19 (20.8%) DENV-3: 4 (4.3%)	median age: 26 (range=6-94)	1.5:1	(Khan et al. 2008)
39	clinically suspected patients, Jeddah	ELISA		Range: 2-60; mean 27.6 $\pm$ 11.2	3.3.:1	(Ayyub et al. 2006)
25	clinically suspected patients, Makkah	RT-PCR	DENV-1 50% DENV-2 33.3%; DENV-3: 16.6%			(Organji et al. 2017)
159	clinically suspected patients, Makkah	ELISA or RT-PCR		25.6 $\pm$ 16.1 years (range 4 to 81 years)	2:1	(Shahin et al. 2009)

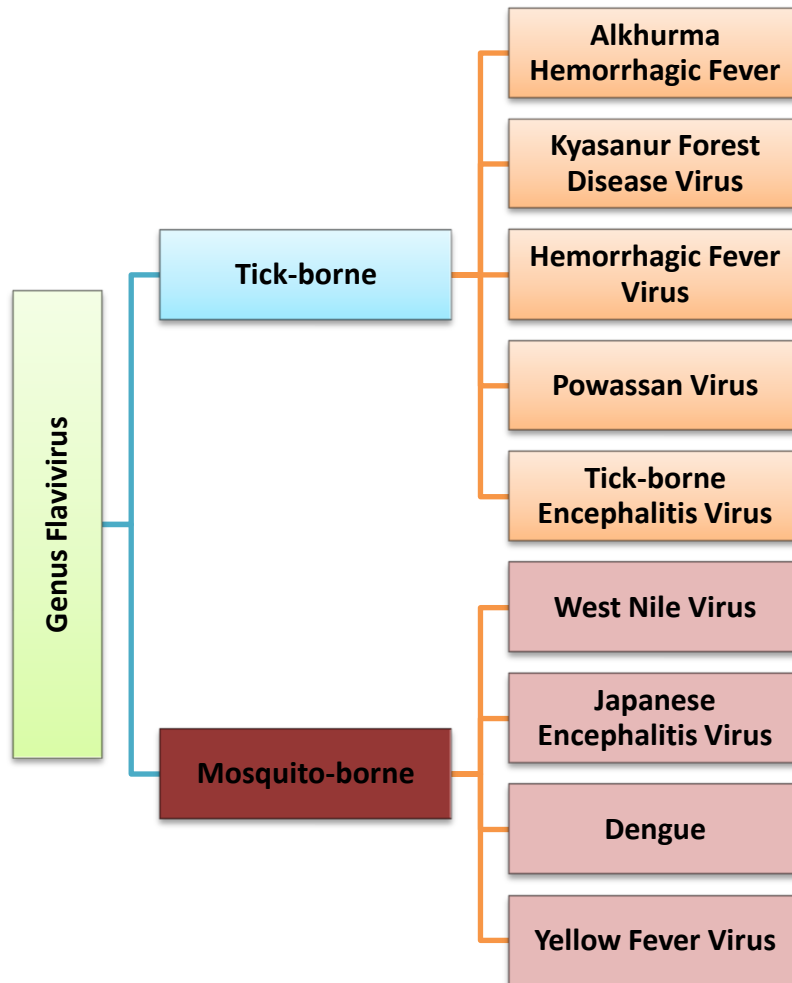
319	clinically suspected patients, Jeddah	Virus culture or serology	DENV-1: 27% DENV-2: 66% DENV-2: 6%	adults between 15-40 years	2.6:1	(Fakeeh & Zaki 2003)
71	admitted to hospitals in Holly Makkah City, 2006 (during the Hajj)	Virus culture or serology		Adults 16-44 years	1.7:1	(El-Gilany et al. 2010)
4187	Confirmed cases reported to the Ministry of Health	NA	NA	47% between 25 and 44 years	2.6:1	(alwafi et al. 2013)
264	Confirmed cases, Jazan	ELISA	NA	52% between 15 and 44 years		(Gamil et al. 2014)
19	Isolates in Jeddah	RT-PCR	DENV-1, DENV-2 and			(Zaki et al. 2008)

			DENV-3			
567	Cross sectional of confirmed cases	RT-PCR in 29%	Not reported	85% were adults	2:1	(Badreddine et al. 2017)

**Table 4: Signs and Symptoms of Patients with Dengue Fever, Numbers are percentage unless indicated otherwise**

	(Khan et al. 2008)	(Ayyub et al. 2006)	(Shahin et al. 2009)	(El-Gilany et al. 2010)
<b>Total number of patients</b>	91	39	159	71
<b>Fever</b>	100	100	100	100
<b>Malaise</b>	83	66.7		67
<b>Musculoskeletal</b>	81		100	59
<b>Headache</b>	75	48.7	100	74
<b>Nausea</b>	69	25.6	27	42.3
<b>Vomiting</b>	65		27	39.4
<b>Abdominal pain</b>	48		24.5	39.4
<b>Dengue fever</b>	93		90	60.5
<b>DHF</b>	7	5	10	39.4
<b>DSS</b>	1		0.6	
<b>Mortality</b>		0	0.6	1.4

**Figure 1: Genus Flavivirus and the Dengue Virus**





**Figure 2: Annual Number of Dengue Fever Cases in Saudi Arabia, Data from** (Alshamrani et al. n.d.; Organji et al. 2017; Saudi Ministry of Health n.d.)

